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Sequence Listing could not be accepted.
If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).
Reviewer: Anne Corrigan
Timestamp: [year=2008; month=8; day=4; hr=13; min=20; sec=52; ms=3;]

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Reviewer Comments:
<210> 1
<211> 217
<212> DNA
<213> Saccharomyces cerevisiae complete genome

The above <213> response is invalid, per 1.823 of the Sequence Rules. The only valid responses are: the Genus species (Genus species only--move other words to the <220>-<223> section. This error appears in many subsequent sequences.

<210> 20
<211> 36
<212> DNA
<213> Sequence Recognized by Synthetic DNA Binding Protein

The above <213> response is invalid, per 1.823 of the Sequence Rules. Please refer to error explanation above for valid <213> responses. Same type of error in Sequences 23, 26-27, 30-34.

****

Validated By CRFValidator v 1.0.3

Application No: 10609383 Version No: 5.0

Input Set:

Output Set:

Started: 2008-08-04 11:53:26.021
Finished: 2008-08-04 11:53:28.365
Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 344 ms
Total Warnings: 34
Total Errors: 0
No. of SeqIDs Defined: 34
Actual SeqID Count: 34

Error code		Error Description
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W	402	Undefined organism found in <213> in SEQ ID (2)
W	402	Undefined organism found in <213> in SEQ ID (3)
W	402	Undefined organism found in <213> in SEQ ID (4)
W	402	Undefined organism found in <213> in SEQ ID (5)
W	402	Undefined organism found in <213> in SEQ ID (6)
W	402	Undefined organism found in <213> in SEQ ID (7)
W	402	Undefined organism found in <213> in SEQ ID (8)
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W	402	Undefined organism found in <213> in SEQ ID (18)
W	402	Undefined organism found in <213> in SEQ ID (19)
W	402	Undefined organism found in <213> in SEQ ID (20)

Input Set:

Output Set:

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Error code	Error Description

This error has occured more than 20 times, will not be displayed

SEQUENCE LISTING

<110> Feldmann, Richard J.

<120> Modifying the Control of Gene Expression Behavior by the Deletion of Connectrons and by the Design and Addition of Synthetic Connectrons in Prokaryotic, Archea and Eukaryotic Genomes

<130> FELD3002CIP1/ESS

<140> 10609383
<141> 2003-07-01

<150> US 09/866,925
<151> 2001-05-30

<150> US 60/393,558
<151> 2002-07-05

<160> 34

<170> PatentIn version 3.5

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<212> DNA
<213> Saccharomyces cerevisiae complete genome

<220>
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<222> (12572)..(12788)
<223> Chromosome = 1 Strand = positive Connectron Object Number = 36

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ggtggtaatg atgaagtaat ttcctgactt gttgttgtac tggtaacagg tggtaatgaa 120

gaagtaattt cctgacttgt tgttgcactg gtaacaggtg gtaatgatga agtaatttcc 180

tgacttgttg ttgtactggt aacaggtggt aatgatg 217

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<211> 236
<212> DNA
<213> Saccharomyces cerevisiae complete genome

<220>
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<222> (12572)..(12807)
<223> Chromosome = 1 Strand = positive Connectron Object Number = 39

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gaagtaattt cctgacttgt tgttgcactg gtaacaggtg gtaatgatga agtaatttcc 180

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<210> 3
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<213> Saccharomyces cerevisiae complete genome

<220>
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<222> (24863)..(25028)
<223> Chromosome = 1 Strand = negative Connectron Object Number = 112

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gtcaggaaat tacttcttca ttaccacctg ttaccactac aaaaacgagc gaacaaacca 120

ctttggttac cgtgacatcc tgcgaatctc atgtgtgcac tgaatc 166

<210> 4
<211> 37
<212> DNA
<213> Escherichia coli k-12 MG1655 complete genome

<220>
<221> misc_feature
<222> (4626130)..(4626166)
<223> Chromosome = 1 Strand = positive Connectron Object Number = 4651a

<400> 4
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<210> 5
<211> 54
<212> DNA
<213> Escherichia coli k12 MG1655 complete genome

<220>
<221> misc_feature
<222> (705150)..(705203)
<223> Chromosome = 1 Strand = negative Connectron Object Number = 811a

<400> 5

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<210> 6
<211> 36
<212> DNA
<213> Escherichia coli k12 MG1655 complete genome

<220>
<221> misc_feature
<222> (757718)..(757753)
<223> Chromosome = 1 Strand = negative Connectron Object Number = 975

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<210> 7
<211> 37
<212> DNA
<213> Escherichia coli k12 MG1655 complete genome

<220>
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<222> (4626130)..(4626166)
<223> Chromosome = 1 Strand = positive Connectron Object Number = 4651a

<400> 7
tctgatgaca aacgccaaac tgcctgatgc gctacgc 37

<210> 8
<211> 54
<212> DNA
<213> Escherichia coli k12 MG1655 complete genome

<220>
<221> misc_feature
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<223> Chromosome = 1 Strand = negative Connectron Object Number = 809

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<210> 9
<211> 36
<212> DNA
<213> Escherichia coli k12 MG1655 complete genome

<220>
<221> misc_feature

<222> (757718)..(757753)
<223> Chromosome = 1 Strand = negtive Connectron Object Number = 975

<400> 9
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<210> 10
<211> 16
<212> DNA
<213> Saccharomyces cerevisiae complete genome - problem

<220>
<221> misc_feature
<222> (221330)..(221345)
<223> Chromosome = 2 Strand = positive Connectron Object Number = 792a

<400> 10
tatatatatg tcactg 16

<210> 11
<211> 16
<212> DNA
<213> Saccharomyces cerevisiae complete genome - problem

<220>
<221> misc_feature
<222> (221346)..(221361)
<223> Chromosome = 2 Strand = positive Connectron Object Number = 793

<400> 11
tattgcatgc tggatg 16

<210> 12
<211> 539
<212> DNA
<213> Saccharomyces cerevisiae complete genome - problem

<220>
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<222> (448454)..(448992)
<223> Chromosome = 5 Strand = positive Connectron Object Number = 4749

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gcatctagga agtaaccttg tacgaaaata ggcaatattt cctgtttagg cgattgtgac 120

gcagatttta gtccaacgat ctagcgtcaa ggaatttttt tatagtggga cattgcacca 180

aggaagtaac ttgatacgtc gtgggtgaat gggtctgttt tcttattcgg cggggtaata 240

catttttggg ggaagtttgt ctgtctgacg cgccatatgt aggtacgcca aaaagggctc 300

ctctacttcg aagcgcgagg tcgtatacct aataaggaaa tgtaatttat aacttttttat 360

tatattggtc ttttcgagag cggaacgtag gtccatgttt aaagtatcca agagaatatc 420

cacgaagcgg ctgagcaacg aacagaatcc tggttctcct cgactaagca gatagttaag 480

atactgtgca ccatggaaat tgaaaacgaa agtacgtacc gactacttta tttttgcag 539

<210> 13
<211> 158
<212> DNA
<213> Saccharomyces cerevisiae complete genome - problem

<220>
<221> misc_feature
<222> (24863)..(25028)
<223> Chromosome = 5 Strand = negative Connectron Object Number = 4824a

<400> 13
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gcatctagga agtaaccttg tacgaaaata ggcaatattt cctgtttagg cgattgtgac 120

gcagatttta gtccaacgat ctagcgtcaa ggaatttt 158

<210> 14
<211> 134
<212> DNA
<213> Halobacterium sp. NRC-1 complete genome

<220>
<221> misc_feature
<222> (732401)..(732534)
<223> Chromosome = 1 Strand = positive Connectron Object Number = 6612

<400> 14
ttcatcacag acgaggacga gcgcggccaa gtggggatcg gcacactcat cgtgttcatc 60

gcgatggtgc tggtcgccgc gatcgccgcc ggcgtcctca tcaacactgc cggctacctc 120

caatccaagg ggtc 134

<210> 15
<211> 193
<212> DNA
<213> Halobacterium sp. NAC-1 complete genome

<220>
<221> misc_feature
<222> (733018)..(733209)
<223> Chromosome = 1 Strand = positive Connectron Object Number = 6644a

<400> 15
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gccgcgatcg ccgccggcgt cctcatcaac accgccggct acctccaatc caaggggtcg 120

gcaaccggtg aggaagcctc cgcacaggtc tccaaccgca tcaacatcgt ctccgcgtac 180

ggcaacgtca aca 193

<210> 16
<211> 85
<212> DNA
<213> Halobacterium sp. NAC-1 complete genome

<220>
<221> misc_feature
<222> (773399)..(773483)
<223> Chromosome = 1 Strand = positive Connectron Object Number = 6852

<400> 16
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ggcgtcctca tcaacactgc cggct 85

<210> 17
<211> 121
<212> DNA
<213> Pseudomonas aeruginosa PA01, complete genome

<220>
<221> misc_feature
<222> (4832718)..(4832838)
<223> Chromosome = 1 Strand = positive Connectron Object Number = 53464

<400> 17
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aagggcatct ccgagcagac caacctgctc gccctcaacg ccgccatcga agccgcgcgc 120

g 121

<210> 18

<211> 194
<212> DNA
<213> Pseudomonas aeruginosa PA01, complete genome

<220>
<221> misc_feature
<222> (4836528)..(4836720)
<223> Chromosome = 1 Strand = positive Connectron Object Number = 53531

<400> 18
ggacggcaaa caggtggtcg agcagaccat ccgcgcgatg aacgagcttt ccgagaagat 60

cagcgcctcc tgcgccaaca tcgaggccct caacagccgc acggtgaaca tcggccagat 120

cctcgaagtg atcaagggca tctccgagca gaccaacctg ctcgccctca acgccgccat 180

cgaagccgcg cgcg 194

<210> 19
<211> 169
<212> DNA
<213> Pseudomonas aeruginosa PA01, complete genome

<220>
<221> misc_feature
<222> (4838678)..(4838846)
<223> Chromosome = 1 Strand = positive Connectron Object Number = 53549a

<400> 19
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gccctcaaca gccgcacggt gaacatcggc cagatcctcg aagtgatcaa gggcatctcc 120

gagcagacca acctgctcgc cctcaacgcc gccatcgaag ccgcgcgcg 169

<210> 20
<211> 36
<212> DNA
<213> Sequence Recognized by Synthetic DNA Binding Protein

<400> 20
tccccatgag catagatatg caggtaggcg gcaagt 36

<210> 21
<211> 136
<212> DNA
<213> Vibrio cholerae chromosome I, complete chromosome

<220>

<221> misc_feature
<222> (952641)..(952777)
<223> Chromosome = 1 Strand = negative Connectron Object Number = 607

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catagataga ctatgtgatt ggggtgaacg aacgtagcca acaccgctgc agcttcaagt 120

aggaagggta tacctt 136

<210> 22
<211> 117
<212> DNA
<213> Vibrio cholerae chromosome I, complete chromosome

<220>
<221> misc_feature
<222> (1005810)..(1005926)
<223> Chromosome = 1 Strand = negative Connectron Object Number = 646

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cagactatgt gattggggtg aacgaacgta gccaataccg ctgcagcttc aagtagg 117

<210> 23
<211> 36
<212> DNA
<213> Sequence Recognized by Synthetic PNA

<400> 23
tccccatgag catagatatg caggtaggcg gcaagt 36

<210> 24
<211> 136
<212> DNA
<213> Vibrio cholerae chromosome I, complete chromosome

<220>
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<222> (952641)..(952777)
<223> Chromosome = 1 Strand = negative Connectron Object Number = 607

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catagataga ctatgtgatt ggggtgaacg aacgtagcca acaccgctgc agcttcaagt 120

aggaagggta tacctt 136

<210> 25
<211> 117
<212> DNA
<213> Vibrio cholerae chromosome I, complete chromosome

<220>
<221> misc_feature
<222> (1005810)..(1005926)
<223> Chromosome = 1 Strand = negative Connectron Object Number = 646

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<211> 15
<212> DNA
<213> Sequence Recognized by Synthetic Linked Pair of DNA Binding Objects

<400> 26
cccgacacaa cctgc 15

<210> 27
<211> 15
<212> DNA
<213> Sequence Recognized by Synthetic Linked Pair of DNA Binding Objects

<400> 27
cccggggttc ccgag 15

<210> 28
<211> 64
<212> DNA
<213> Aeropyrum pernix k1 complete genome

<220>
<221> misc_feature
<222> (284008)..(284070)
<223> Chromosome = 1 Strand = negative Connectron Object Number = 218

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cggc 64

<210> 29
<211> 163
<212> DNA
<213> Aeropyrum pernix k1 complete genome

<220>
<221> misc_feature
<222> (326716)..(326878)
<223> Chromosome = 1 Strand = negative Connectron Object Number = 295

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ggccccgggg agaccgtgat gaacccagcc gtgcccgaca caacctgcta taatttgtta 120

catgaaggca cggtttgggt gaacggctca taatcctctc gat 163

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<212> DNA
<213> Synthetic Sequence

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<210> 31
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<400> 31
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<213> Synthetic Sequence

<400> 32
uagaggagua ccac 14

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<400> 33
gugguacucc ucua 14

<210> 34
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